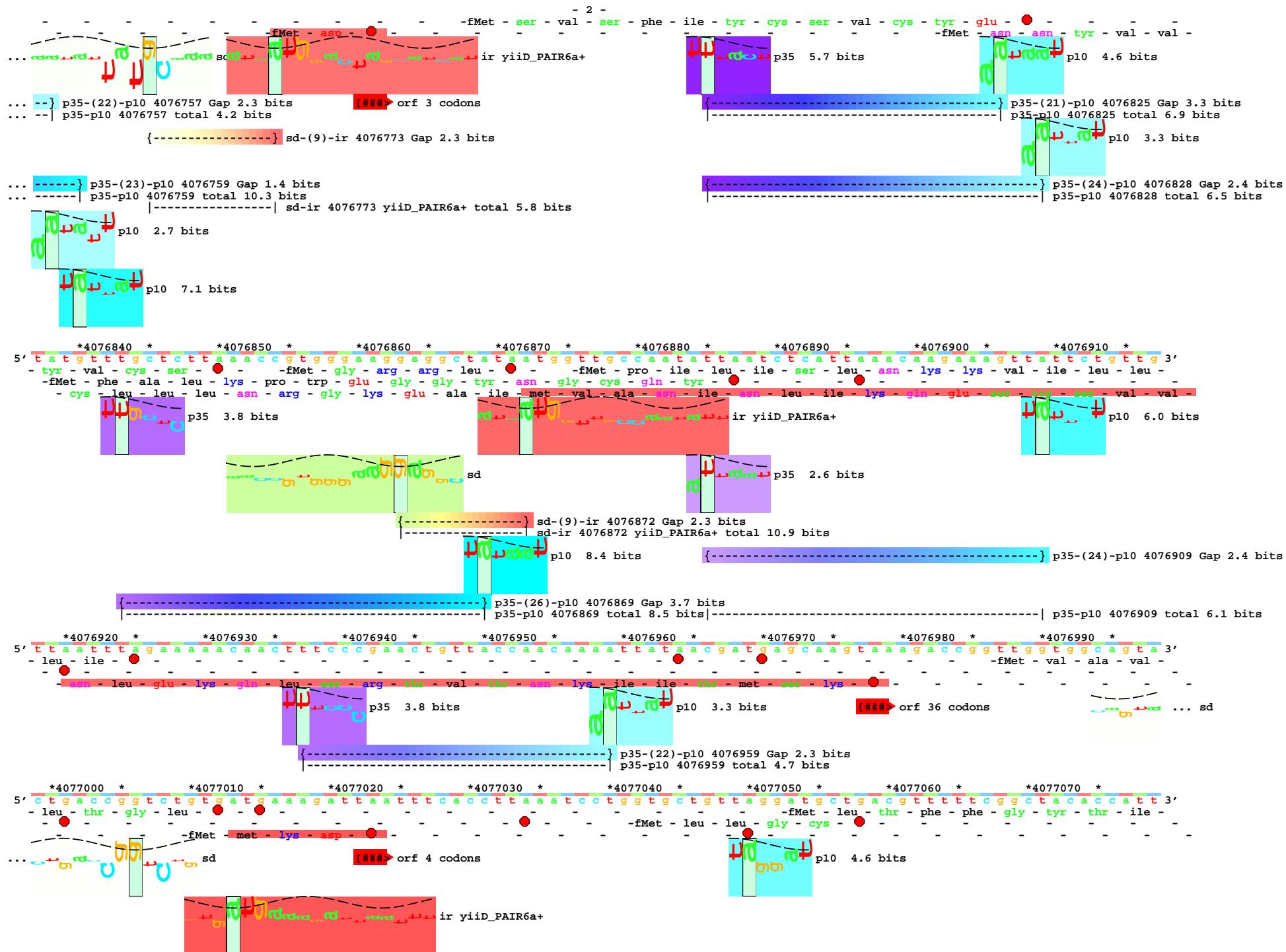


- 1 -
piece 1, NC_000913, yiID PAIR6a+, config: linear, direction: +, begin: 4076432, end: 4077218





- 3 -

```
{-----} sd-(7)-ir 4077013 Gap 3.7 bits  
{-----} sd-ir 4077013 yiiD_PAIR6a+ total 5.6 bits
```

 p35 3.8 bits

```
{- } p35-(24)-p10 4077050 Gap 2.4 bits  
|-----| p35-p10 4077050 total 6.0 bits
```

{-----} p35-(21)-p10 4077108 Gap 3.3 bits

|-----| r35-p10 4077108 total 8.1 bits

{-----} p35-(23)-p10 4077110 Gap 1.4 bits
{-----} p35-p10 4077110 total 7.8 bits

+10 +10 +10 p10 6.5 bits

{-----} p35-(26)-p10 4077113 Gap 3.7 bits
{-----} p35-p10 4077113 total 5.8 bits

The diagram illustrates the transcription of the *yiid_PPAIR6a+* gene. The top part shows the DNA template strand with transcription start sites marked by asterisks (*). The bottom part shows the primary transcript with the deduced amino acid sequence. The sequence starts with fMet and ends with lys. The mature mRNA is shown in red, while the 5' UTR and 3' UTR are in black. The mature mRNA contains a poly-A tail (aa...aa) at the 3' end.

{-----} sd-(9)-ir 4077171 Gap 2.3 bits

p10 7.1 bits

```
| sd-ir 4077171 viID PAIR6a+ total 7.1 bits  
{-----} p35-(24)-p10 4077185 Gap 2.4 bits  
|-----| p35-p10 4077185 total 4.3 bits
```

p35 3.3 bits

{-----} p35-(21)-p10 4077193 Gap 3.3 bits
----- p35-p10 4077193 total 7.1 bits